

RAW SEQUENCE LISTING

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Application Serial Number: 10/705,401
Source: 1 FWO
Date Processed by STIC: 10/18/04

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RAW SEQUENCE LISTING

DATE: 10/18/2004

PATENT APPLICATION: US/10/705,401

TIME: 14:36:20

Input Set : A:\PTO.FG.txt

Output Set: N:\CRF4\10182004\J705401.raw

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3 <110> APPLICANT: Clausen, Henrik
4   Bennett, Eric P.
6 <120> TITLE OF INVENTION: METHODS TO IDENTIFY AGENTS MODULATING FUNCTIONS OF
POLYPEPTIDE GALNAC-
7   TRANSFERASES, PHARMACEUTICAL COMPOSITIONS COMPRISING SUCH AGENTS AND THE
8   USE OF SUCH AGENTS FOR PREPARING MEDICAMENTS
W--> 9 <130> FILE REFERENCE: 04305/100H154-US2
C--> 11 <140> CURRENT APPLICATION NUMBER: US/10/705,401
C--> 11 <141> CURRENT FILING DATE: 2003-11-10
11 <150> PRIOR APPLICATION NUMBER: US 60/425,204
12 <151> PRIOR FILING DATE: 2002-11-08
14 <150> PRIOR APPLICATION NUMBER: PCT/DK03/00763
15 <151> PRIOR FILING DATE: 2003-11-07
17 <160> NUMBER OF SEQ ID NOS: 127
19 <170> SOFTWARE: PatentIn version 3.1
21 <210> SEQ ID NO: 1
22 <211> LENGTH: 20
23 <212> TYPE: PRT
24 <213> ORGANISM: Artificial Sequence
26 <220> FEATURE:
27 <223> OTHER INFORMATION: synthetic peptide
29 <400> SEQUENCE: 1
31 His Gly Val Thr Ser Ala Pro Asp Thr Arg Pro Ala Pro Gly Ser Thr
32 1           5           10           15
35 Ala Pro Pro Ala
36           20
39 <210> SEQ ID NO: 2
40 <211> LENGTH: 24
41 <212> TYPE: PRT
42 <213> ORGANISM: Artificial Sequence
44 <220> FEATURE:
45 <223> OTHER INFORMATION: synthetic peptide
47 <400> SEQUENCE: 2
49 Thr Ala Pro Pro Ala His Gly Val Thr Ser Ala Pro Asp Thr Arg Pro
50 1           5           10           15
53 Ala Pro Gly Ser Thr Ala Pro Pro
54           20
57 <210> SEQ ID NO: 3
58 <211> LENGTH: 167
59 <212> TYPE: PRT
60 <213> ORGANISM: Homo sapiens
62 <400> SEQUENCE: 3
64 Tyr Gly Asp Ile Ser Ser Arg Val Gly Leu Arg His Lys Leu Gln Cys
65 1           5           10           15

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68 Lys Pro Phe Ser Trp Tyr Leu Glu Asn Ile Tyr Pro Asp Ser Gln Ile
69          20          25          30
72 Pro Arg His Tyr Phe Ser Leu Gly Glu Ile Arg Asn Val Glu Thr Asn
73          35          40          45
76 Gln Cys Leu Asp Asn Met Ala Arg Lys Glu Asn Glu Lys Val Gly Ile
77          50          55          60
80 Phe Asn Cys His Gly Met Gly Gly Asn Gln Val Phe Ser Tyr Thr Ala
81 65          70          75          80
84 Asn Lys Glu Ile Arg Thr Asp Asp Leu Cys Leu Asp Val Ser Lys Leu
85          85          90          95
88 Asn Gly Pro Val Thr Met Leu Lys Cys His His Leu Lys Gly Asn Gln
89          100          105          110
92 Leu Trp Glu Tyr Asp Pro Val Lys Leu Thr Leu Gln His Val Asn Ser
93          115          120          125
96 Asn Gln Cys Leu Asp Lys Ala Thr Glu Glu Asp Ser Gln Val Pro Ser
97          130          135          140
100 Ile Arg Asp Cys Asn Gly Ser Arg Ser Gln Gln Trp Leu Leu Arg Asn
101 145          150          155          160
104 Val Thr Leu Pro Glu Ile Phe
105          165
108 <210> SEQ ID NO: 4
109 <211> LENGTH: 164
110 <212> TYPE: PRT
111 <213> ORGANISM: Homo sapiens
113 <400> SEQUENCE: 4
115 Tyr Gly Asn Ile Gln Ser Arg Leu Glu Leu Arg Lys Lys Leu Ser Cys
116 1          5          10          15
119 Lys Pro Phe Lys Trp Tyr Leu Glu Asn Val Tyr Pro Glu Leu Arg Val
120          20          25          30
123 Pro Asp His Gln Asp Ile Ala Phe Gly Ala Leu Gln Gln Gly Thr Asn
124          35          40          45
127 Cys Leu Asp Thr Leu Gly His Phe Ala Asp Gly Val Val Gly Val Tyr
128          50          55          60
131 Glu Cys His Asn Ala Gly Gly Asn Gln Glu Trp Ala Leu Thr Lys Glu
132 65          70          75          80
135 Lys Ser Val Lys His Met Asp Leu Cys Leu Thr Val Val Asp Arg Ala
136          85          90          95
139 Pro Gly Ser Leu Ile Lys Leu Gln Gly Cys Arg Glu Asn Asp Ser Arg
140          100          105          110
143 Gln Lys Trp Glu Gln Ile Glu Gly Asn Ser Lys Leu Arg His Val Gly
144          115          120          125
147 Ser Asn Leu Cys Leu Asp Ser Arg Thr Ala Lys Ser Gly Gly Leu Ser
148          130          135          140
151 Val Glu Val Cys Gly Pro Ala Leu Ser Gln Gln Trp Lys Phe Thr Leu
152 145          150          155          160
155 Asn Leu Gln Gln
159 <210> SEQ ID NO: 5
160 <211> LENGTH: 167
161 <212> TYPE: PRT

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162 <213> ORGANISM: Homo sapiens
164 <400> SEQUENCE: 5
166 Phe Gly Asp Leu Ser Lys Arg Phe Glu Ile Lys His Arg Leu Arg Cys
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170 Lys Asn Phe Thr Trp Tyr Leu Asn Asn Ile Tyr Pro Glu Val Tyr Val
171 20 25 30
174 Pro Asp Leu Asn Pro Val Ile Ser Gly Tyr Ile Lys Ser Val Gly Gln
175 35 40 45
178 Pro Leu Cys Leu Asp Val Gly Glu Asn Asn Gln Gly Gly Lys Pro Leu
179 50 55 60
182 Ile Met Tyr Thr Cys His Gly Leu Gly Gly Asn Gln Tyr Phe Glu Tyr
183 65 70 75 80
186 Ser Ala Gln His Glu Ile Arg His Asn Ile Gln Lys Glu Leu Cys Leu
187 85 90 95
190 His Ala Ala Gln Gly Leu Val Gln Leu Lys Ala Cys Thr Tyr Lys Gly
191 100 105 110
194 His Lys Thr Val Val Thr Gly Glu Gln Ile Trp Glu Ile Gln Lys Asp
195 115 120 125
198 Gln Leu Leu Tyr Asn Pro Phe Leu Lys Met Cys Leu Ser Ala Asn Gly
199 130 135 140
202 Glu His Pro Ser Leu Val Ser Cys Asn Pro Ser Asp Pro Leu Gln Lys
203 145 150 155 160
206 Trp Ile Leu Ser Gln Asn Asp
207 165
210 <210> SEQ ID NO: 6
211 <211> LENGTH: 174
212 <212> TYPE: PRT
213 <213> ORGANISM: Homo sapiens
215 <400> SEQUENCE: 6
217 Ala Tyr Gly Asp Ile Ser Glu Arg Lys Leu Leu Arg Glu Arg Leu Arg
218 1 5 10 15
221 Cys Lys Ser Phe Asp Trp Tyr Leu Lys Asn Val Phe Pro Asn Leu His
222 20 25 30
225 Val Pro Glu Asp Arg Pro Gly Trp His Gly Ala Ile Arg Ser Arg Gly
226 35 40 45
229 Ile Ser Ser Glu Cys Leu Asp Tyr Asn Ser Pro Asp Asn Asn Pro Thr
230 50 55 60
233 Gly Ala Asn Leu Ser Leu Phe Gly Cys His Gly Gln Gly Gly Asn Gln
234 65 70 75 80
237 Phe Phe Glu Tyr Thr Ser Asn Lys Glu Ile Arg Phe Asn Ser Val Thr
238 85 90 95
241 Glu Leu Cys Ala Glu Val Pro Glu Gln Lys Asn Tyr Val Gly Met Gln
242 100 105 110
245 Asn Cys Pro Lys Asp Gly Phe Pro Val Pro Ala Asn Ile Ile Trp His
246 115 120 125
249 Phe Lys Glu Asp Gly Thr Ile Phe His Pro His Ser Gly Leu Cys Leu
250 130 135 140
253 Ser Ala Tyr Arg Thr Pro Glu Gly Arg Pro Asp Val Gln Met Arg Thr
254 145 150 155 160

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257 Cys Asp Ala Leu Asp Lys Asn Gln Ile Trp Ser Phe Glu Lys
258             165             170
261 <210> SEQ ID NO: 7
262 <211> LENGTH: 168
263 <212> TYPE: PRT
264 <213> ORGANISM: Homo sapiens
266 <400> SEQUENCE: 7
268 Asp Val Gly Asn Leu Thr Gln Gln Arg Glu Leu Arg Lys Lys Leu Lys
269 1             5             10             15
272 Cys Lys Ser Phe Lys Trp Tyr Leu Glu Asn Val Phe Pro Asp Leu Arg
273             20             25             30
276 Ala Pro Ile Val Arg Ala Ser Gly Val Leu Ile Asn Val Ala Leu Gly
277             35             40             45
280 Lys Cys Ile Ser Ile Glu Asn Thr Thr Val Ile Leu Glu Asp Cys Asp
281             50             55             60
284 Gly Ser Lys Glu Leu Gln Gln Phe Asn Tyr Thr Trp Leu Arg Leu Ile
285 65             70             75             80
288 Lys Cys Gly Glu Trp Cys Ile Ala Pro Ile Pro Asp Lys Gly Ala Val
289             85             90             95
292 Arg Leu His Pro Cys Asp Asn Arg Asn Lys Gly Leu Lys Trp Leu His
293             100            105            110
296 Lys Ser Thr Ser Val Phe His Pro Glu Leu Val Asn His Ile Val Phe
297             115            120            125
300 Glu Asn Asn Gln Gln Leu Leu Cys Leu Glu Gly Asn Phe Ser Gln Lys
301             130            135            140
304 Ile Leu Lys Val Ala Ala Cys Asp Pro Val Lys Pro Tyr Gln Lys Trp
305 145            150            155            160
308 Lys Phe Glu Lys Tyr Tyr Glu Ala
309             165
312 <210> SEQ ID NO: 8
313 <211> LENGTH: 165
314 <212> TYPE: PRT
315 <213> ORGANISM: Homo sapiens
317 <400> SEQUENCE: 8
319 Ser Phe Gly Asp Ile Ser Glu Arg Leu Gln Leu Arg Glu Gln Leu His
320 1             5             10             15
323 Cys His Asn Phe Ser Trp Tyr Leu His Asn Val Tyr Pro Glu Met Phe
324             20             25             30
327 Val Pro Asp Leu Thr Pro Thr Phe Tyr Gly Ala Ile Lys Asn Leu Gly
328             35             40             45
331 Thr Asn Gln Cys Leu Asp Val Gly Glu Asn Asn Arg Gly Gly Lys Pro
332             50             55             60
335 Leu Ile Met Tyr Ser Cys His Gly Leu Gly Gly Asn Gln Tyr Phe Glu
336 65             70             75             80
339 Tyr Thr Thr Gln Arg Asp Leu Arg His Asn Ile Ala Lys Gln Leu Cys
340             85             90             95
343 Leu His Val Ser Lys Gly Ala Leu Gly Leu Gly Ser Cys His Phe Thr
344             100            105            110
347 Gly Lys Asn Ser Gln Val Pro Lys Asp Glu Glu Trp Glu Leu Ala Gln

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348          115          120          125
351 Asp Gln Leu Ile Arg Asn Ser Gly Ser Gly Thr Cys Leu Thr Ser Gln
352          130          135          140
355 Asp Lys Lys Pro Ala Met Ala Pro Cys Asn Pro Ser Asp Pro His Gln
356 145          150          155          160
359 Leu Trp Leu Phe Val
360          165
363 <210> SEQ ID NO: 9
364 <211> LENGTH: 166
365 <212> TYPE: PRT
366 <213> ORGANISM: Homo sapiens
368 <400> SEQUENCE: 9
370 Tyr Gly Asp Ile Ser Glu Leu Lys Lys Phe Arg Glu Asp His Asn Cys
371 1          5          10          15
374 Gln Ser Phe Lys Trp Phe Met Glu Glu Ile Ala Tyr Asp Ile Thr Ser
375          20          25          30
378 His Tyr Pro Leu Pro Pro Lys Asn Val Asp Trp Gly Glu Ile Arg Gly
379          35          40          45
382 Phe Glu Thr Ala Tyr Cys Ile Asp Ser Met Gly Lys Thr Asn Gly Gly
383          50          55          60
386 Phe Val Glu Leu Gly Pro Cys His Arg Met Gly Gly Asn Gln Leu Phe
387 65          70          75          80
390 Arg Ile Asn Glu Ala Asn Gln Leu Met Gln Tyr Asp Gln Cys Leu Thr
391          85          90          95
394 Lys Gly Ala Asp Gly Ser Lys Val Met Ile Thr His Cys Asn Leu Asn
395          100          105          110
398 Glu Phe Lys Glu Trp Gln Tyr Phe Lys Asn Leu His Arg Phe Thr His
399          115          120          125
402 Ile Pro Ser Gly Lys Cys Leu Asp Arg Ser Glu Val Leu His Gln Val
403          130          135          140
406 Phe Ile Ser Asn Cys Asp Ser Ser Lys Thr Thr Gln Lys Trp Glu Met
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410 Asn Asn Ile His Ser Val
411          165
414 <210> SEQ ID NO: 10
415 <211> LENGTH: 179
416 <212> TYPE: PRT
417 <213> ORGANISM: Homo sapiens
419 <400> SEQUENCE: 10
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425 Lys Thr Phe Asp Trp Tyr Leu Lys Asn Val Tyr Pro Leu Leu Lys Pro
426          20          25          30
429 Leu His Thr Ile Val Gly Tyr Gly Arg Met Lys Asn Leu Leu Asp Glu
430          35          40          45
433 Asn Val Cys Leu Asp Gln Gly Pro Val Pro Gly Asn Thr Pro Ile Met
434          50          55          60
437 Tyr Tyr Cys His Glu Phe Ser Ser Gln Asn Val Tyr Tyr His Leu Thr
438 65          70          75          80

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RAW SEQUENCE LISTING ERROR SUMMARY

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Input Set : A:\PTO.FG.txt

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Invalid Line Length:

The rules require that a line not exceed 72 characters in length. This includes spaces.

Seq#:1; Line(s) 7

VERIFICATION SUMMARY

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DATE: 10/18/2004

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Input Set : A:\PTO.FG.txt

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L:9 M:283 W: Missing Blank Line separator, <130> field identifier
L:11 M:270 C: Current Application Number differs, Replaced Current Application No
L:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date